

#5 1/270
4/24/01
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OIFE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/685,696
DATE: 10/19/2000
TIME: 15:06:07

Input Set : A:\455c13.app
Output Set : N:\CRF3\10192000\I685696.raw

P.5

ENTERED

```
3 <110> APPLICANT: Wang, Tongtong
4   Fan, Liqun
5   Kalos, Michael D.
6   Bangur, Chaitanya S.
7   Hosken, Nancy
8   Fanger, Gary R.
9   Li, Samuel X.
10  Wang, Aijun
11  Skeiky, Yasir A.W.
12  Henderson, Robert A.
13  McNeill, Patricia D.
18 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
19   AND DIAGNOSIS OF LUNG CANCER
21 <130> FILE REFERENCE: 210121.455C13
C--> 23 <140> CURRENT APPLICATION NUMBER: US/09/685,696
24 <141> CURRENT FILING DATE: 2000-10-09
26 <160> NUMBER OF SEQ ID NOS: 381
28 <170> SOFTWARE: FastSEQ for Windows Version 3.0
30 <210> SEQ ID NO: 1
31 <211> LENGTH: 315
32 <212> TYPE: DNA
33 <213> ORGANISM: Homo sapien
35 <220> FEATURE:
36 <221> NAME/KEY: misc_feature
37 <222> LOCATION: (1)...(315)
38 <223> OTHER INFORMATION: n = A,T,C or G
40 <400> SEQUENCE: 1
41   gcagagacag actggtgggtt gaacctggag gtgccaaaaa agccagctgc gggcccagga      60
42   cagctgccgt gagactcccg atgtcacagg cagtctgtgt ggttacagcg cccctcagtg      120
43   ttcattctcca gcagagacaa cggaggaggc tcccaccagg acggttctca ttatttatat      180
W--> 44   gttaatatgt ttgtaaactc atgtacagtt tttttgggg gggaagcaat gggaanggta      240
W--> 45   naaattacaa atagaatcat ttgctgtaat ccttaaatgg caaacggtea ggccacgtga      300
46   aaaaaaaaaa aaaaaa                                     315
48 <210> SEQ ID NO: 2
49 <211> LENGTH: 380
50 <212> TYPE: DNA
51 <213> ORGANISM: Homo sapien
53 <400> SEQUENCE: 2
54   atttaggctt aagattttgt ttacccttgt tactaaggag caaattagta ttaaagtata      60
55   atatatataa acaaatataa aaagttttga gtggttcagc tttttatatt tttttaatgg      120
56   cataactttt aacaacactg ctctgtaaat ggttgaactg tggtaactag actgagataa      180
57   ctgaaatgag tggatgtata gtgtatttgc ataattatcc cactatgaag caaagggact      240
58   ggataaattc ccagtcctaga ttattagcct ttgttaacca tcaagcacct agaagaagaa      300
59   ttattggaaa ttttgccttc tgtaactggc actttggggg gtgacttate ttttgccttt      360
60   gtaaaaaaaaaa aaaaaaaaaa                                     380
62 <210> SEQ ID NO: 3
63 <211> LENGTH: 346
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64 <212> TYPE: DNA
65 <213> ORGANISM: Homo sapien
67 <220> FEATURE:
68 <221> NAME/KEY: misc_feature
69 <222> LOCATION: (1)...(346)
70 <223> OTHER INFORMATION: n = A,T,C or G
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74 catcctcacc ataccaccatc cactttccaa taacatttaa tcctttctaa aattgtaagt 120
75 atacaattgt actttctttg gattttcata acaaatatac catagactgt taattttatt 180
76 gaagtttcct taatggaatg agtcattttt gtcttggtgt tttgagggtta cctttgcttt 240
77 gacttccaac aatttgatca tatagtgttg agctgtggaa atctttaagt ttattctata 300
W--> 78 gcaataattt ctattnnnag annccngggn naaaannann annaaa 346
80 <210> SEQ ID NO: 4
81 <211> LENGTH: 372
82 <212> TYPE: DNA
83 <213> ORGANISM: Homo sapien
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86 <221> NAME/KEY: misc_feature
87 <222> LOCATION: (1)...(372)
88 <223> OTHER INFORMATION: n = A,T,C or G
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92 tggtttggtt tggttttttg aactggtagt taggggtggt cacagttcta atgtaagcac 120
93 tctcttctcc aagtgtgtgt ttgtggggac aatcattctt tgaacattag agaggaaggg 180
94 agttcaagct gttgaaaaga ctattgctta tttttgtttt taaagacctt cttgacgtca 240
W--> 95 tgtggacagt gcacgtgcct tacgctacat cttgttttct aggaagaagg ggatgcnggg 300
W--> 96 aaggantggg tgctttgtga tggataaaac gnctaaataa cacaccttta cattttgaaa 360
97 aaacaaaaac aa 372
99 <210> SEQ ID NO: 5
100 <211> LENGTH: 698
101 <212> TYPE: DNA
102 <213> ORGANISM: Homo sapien
104 <220> FEATURE:
105 <221> NAME/KEY: misc_feature
106 <222> LOCATION: (1)...(698)
107 <223> OTHER INFORMATION: n = A,T,C or G
109 <400> SEQUENCE: 5
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111 cctaaccacg gtttaactgca agaagaggcg ggatactttc agctttccat gtaactgtat 120
112 gcataaagcc aatgtagtcc agtttctaag atcatgttcc aagctaaactg aatccactt 180
113 caatacacac tcatgaactc ctgatggaac aataacaggg ccaagcctgt ggtatgatgt 240
114 gcacacttgc tagactcaga aaaaatacta ctctcataaa tgggtgggag tattttgggt 300
W--> 115 gacaacctac tttgcttggc tgagtgaagg aatgatattc atatnttcat ttattccatg 360
116 gacatttagt tagtgctttt tatataccag gcatgatgct gaggacact cttgtgtata 420
W--> 117 tntccaaatn ttngtncngt cgctgcacat atctgaaatc ctatattaag antttcccaa 480
W--> 118 natgangtcc ctggtttttc cagccactt gatcngtcaa ngatctcacc tctgntgtgc 540
W--> 119 ctaaaacctn ctncnnnang gttagacnng acctctcttc tcccttcccg aanaatnaag 600
W--> 120 tgtgngaaga nanccnncn cccctctnnc tncnncctng ccngctnnnc cncntgtngg 660

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W--> 121 gggngccgcc cccgcggggg gacccccccn ttttcccc      698
123 <210> SEQ ID NO: 6
124 <211> LENGTH: 740
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126 <213> ORGANISM: Homo sapien
128 <220> FEATURE:
129 <221> NAME/KEY: misc_feature
130 <222> LOCATION: (1)...(740)
131 <223> OTHER INFORMATION: n = A,T,C or G
133 <400> SEQUENCE: 6
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136 gccaatattt ccttatatct atccataaca ttatactac atttgtaaga gaatatgcac      180
137 gtgaaactta acactttata aggtaaaaat gaggtttcca agatttaata atctgatcaa      240
138 gttcttggtt ttcccaataa gaatggactt ggtctgttaa ggggctaagg gagaagaaga      300
139 agataagggtt aaaagtgttt aatgaccaa cttctaaaaa gaaatgcaa aaaaaattta      360
W--> 140 ttttcaagcc ttcgaactat ttaaggaaa caaatcatt tcctanatgc atatcatttg      420
W--> 141 tgagantttc tcantaatat cctgaatcat tcatttcagc tnaggcttca tgttgactcg      480
W--> 142 atatgtcatc tagggaaagt ctatttcatg gtccaaacct gttgccatag ttggttaggc      540
W--> 143 tttcctttaa ntgtgaanta ttacangaa attttctct tnanagtct tnatagggtt      600
W--> 144 aggggtgtgg gaaaagcttc taacaatctg tagtgttncg tgttatctgt ncagaaccan      660
W--> 145 aatnacggat cgnangaagg actgggtcta ttacangaa cgaatnatct ngttnnntgt      720
W--> 146 gtnnncaact ccngggagcc      740
148 <210> SEQ ID NO: 7
149 <211> LENGTH: 670
150 <212> TYPE: DNA
151 <213> ORGANISM: Homo sapien
153 <220> FEATURE:
154 <221> NAME/KEY: misc_feature
155 <222> LOCATION: (1)...(670)
156 <223> OTHER INFORMATION: n = A,T,C or G
158 <400> SEQUENCE: 7
159 gctggggagc tcggcatggc ggtccccgct gcagccatgg ggcctcggc gttgggccag      60
160 agcggccccc gctcgatggc cccgtgggtc tcagtgaaga gcggcccgtc gcgctacgtg      120
161 cttgggatgc aggagctgtt ccggggccac agcaagaccg cgagttcctg gcgcacagcg      180
162 ccaagggtga ctcggtggcc tggagtgcg acgggcgtcg cctacctcgg ggtcttcgac      240
W--> 163 aagacgccac gtcttcttgc tgganaanga ccgttgggtca aagaaaacaa ttatcgggga      300
164 catggggata gtgtggacca ctttgttggc atccaagtaa tcctgacctt tttgttacgg      360
165 cgtctggaga taaaccatt cgcactctgg atgtgaggac tacaaaatgc attgccactg      420
W--> 166 tgaacactaa aggggagaa attaatatct gctggantcc tgatgggcan accattgctg      480
W--> 167 tagcnacaag gatgatgtgg tgactttatt gatgccaaga aaccccggtc caaagcaaaa      540
W--> 168 aaacanttcc aanttcgaag tcaccnaaat ctctggaac aatgaacatn aatatnttct      600
W--> 169 tcctgacaat ggnccctggg tgnatcacat cctcagctnc cccaaaactg aancctgtnc      660
W--> 170 natccacccc      670
172 <210> SEQ ID NO: 8
173 <211> LENGTH: 689
174 <212> TYPE: DNA
175 <213> ORGANISM: Homo sapien
177 <220> FEATURE:

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178 <221> NAME/KEY: misc_feature
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180 <223> OTHER INFORMATION: n = A,T,C or G
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184 aaatgaagta ctggatttgg gaaaacctgg ttttattaga acatatggaa tgaaagccta      120
185 cacctagcat tgccacttta gcccccctgaa ttaacagagc ccaattgaga caaacccctg      180
186 gcaacaggaa attcaaggga gaaaaagtaa gcaacttggg ctaggatgag ctgactccct      240
W--> 187 tagagcaaa ganagacagc cccattacc aaataccatt ttgcctggg gcttgtgcag      300
W--> 188 ctggcagtgt tcctgcccc gcctggcacc ttatngtttt gatagcaact tcgttgaatt      360
W--> 189 ttaccaact tattacttga aattataata tagcctgtcc gtttctgttn tccaggctgt      420
W--> 190 gatatatntt cctagtgtgt tgacttttaa aataaatnag gtttantttt cccccccn      480
W--> 191 cnnctnctncc nntcncctnn cnncccccc cncctngtcc tccnnnttn gggggggcnn      540
W--> 192 cccccnccgn ggacccccct ttggtccctt agtggagggt natggcccc ggnnttatcc      600
W--> 193 nggcctann tttccccgtn nnaaatgntt cccccccca ntccnccac ctcaanccgg      660
W--> 194 aagcctaagt ttntaccctg ggggtccccc      689

196 <210> SEQ ID NO: 9
197 <211> LENGTH: 674
198 <212> TYPE: DNA
199 <213> ORGANISM: Homo sapien
201 <220> FEATURE:
202 <221> NAME/KEY: misc_feature
203 <222> LOCATION: (1)...(674)
204 <223> OTHER INFORMATION: n = A,T,C or G
206 <400> SEQUENCE: 9
207 gtccactctc ctttgagtgt actgtcttac tgtgcactct gtttttcaac ttcttagata      60
208 taaaaaatgc ttgttctata gtggagtaag agctcacaca ccaaggcag caagataact      120
209 gaaaaaagcg aggccttttt gccaccttgg taaaggccag ttcactgcta tagaactgct      180
210 ataagcctga aggggaagtag ctatgagact ttccattttt cttagtcttc ccaataggct      240
211 ctttcatgga aaaaggcttc ctgtaataat ttccacctaa tgaattagca gtgtgattat      300
212 ttctgaaata agagacaaat tgggccgcag agtcttctct tgatttaaaa taaacaaccc      360
213 aaagttttgt ttggtcttca ccaaaggaca tactctaggg ggtatgttgt tgaagacatt      420
214 caaaaacatt agctgttctg tctttcaatt tcaagttatt ttggagactg cctccatgtg      480
215 agttaattac ttgtctctgg aactagcatt attgtcatta tcatcacatt ctgtcatcat      540
216 catctgaata atattgtgga ttccccctc tgcttgcatc ttcttttgac tcctctggga      600
W--> 217 anaaatgtca aaaaaaagg tcgatctact cngcaaggnc catctaata ctgcgctgga      660
W--> 218 aggacccnct gcccc      674

220 <210> SEQ ID NO: 10
221 <211> LENGTH: 346
222 <212> TYPE: DNA
223 <213> ORGANISM: Homo sapien
225 <220> FEATURE:
226 <221> NAME/KEY: misc_feature
227 <222> LOCATION: (1)...(346)
228 <223> OTHER INFORMATION: n = A,T,C or G
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232 ttctgtctgt aacaaaaatg tactttatag agatggagga aaaggtctaa tactacatag      120
233 ccttaagtgt ttctgtcatt gttcaagtgt attttctgta acagaaacat atttggaatg      180

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RAW SEQUENCE LISTING

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Input Set : A:\455c13.app

Output Set: N:\CRF3\10192000\I685696.raw

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234 tttttctttt ccccttataa attgtaattc ctgaaatact gctgctttaa aaagtccac 240
235 tgtcagatta tattatctaa caattgaata ttgtaaatat acttgcttta cctctcaata 300
W--> 236 aaagggtact ttctattan nnagnngnnn gnnnnataaa anaana 346
238 <210> SEQ ID NO: 11
239 <211> LENGTH: 602
240 <212> TYPE: DNA
241 <213> ORGANISM: Homo sapien
243 <400> SEQUENCE: 11
244 actagtaaaa agcagcattg ccaaataatc cctaattttc cactaaaaat ataataaat 60
245 gatgttaagc tttttgaaaa gttaggttta aacctactgt tgtagatta atgtatttgt 120
246 tgcctccctt tatctggaat gtggcattag cttttttatt ttaacctctt ttaattctta 180
247 ttaattcca tgaactaagg ttggagagct aaacactggg atttttggat aacagactga 240
248 cagttttgca taattataat cggcattgta catagaaagg atagggctac cttttgttaa 300
249 atctgcactt tctaataatc aaaaaaggga aatgaagtta taaatcaatt ttgtataat 360
250 ctgtttgaaa catgagtttt atttgcttaa tattagggct ttgccctttt tctgtaagtc 420
251 tcttggggtc ctgtgtagaa ctgttctcat taaacaccaa acagttaagt ccattctctg 480
252 gtactagcta caaattcggg ttcatattct acttaacaat ttaataaac tgaaatattt 540
253 ctataggtgc tacttctgtt catataaaaa caaaacttga ttccaaaaa aaaaaaaaaa 600
254 aa 602
256 <210> SEQ ID NO: 12
257 <211> LENGTH: 685
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259 <213> ORGANISM: Homo sapien
261 <220> FEATURE:
262 <221> NAME/KEY: misc_feature
263 <222> LOCATION: (1)...(685)
264 <223> OTHER INFORMATION: n = A,T,C or G
266 <400> SEQUENCE: 12
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268 attatcatgg tattgatgga cctaagaaaa taaaaattag actaagcccc caaataagct 120
W--> 269 gcatgcattt gtaacatgat tagtagattt gaatatatag atgtagtatn ttgggtatct 180
270 aggtgtttta tcattatgta aaggaattaa agtaaaggac tttgtagttg tttttattaa 240
W--> 271 atatgcatat agtagagtgc aaaaatatag caaaaatana aactaaagggt agaaaagcat 300
W--> 272 tttagatatg ccttaatnta nnaactgtgc caggtggccc tcggaataga tgccaggcag 360
273 agaccagtgc ctgggtgggt cctccccttg tctgcccccc tgaagaactt ccctcacgtg 420
W--> 274 angtagtgcc ctgtaggtg tcacgtggan tantggganc aggccgnncn gtnanaagaa 480
W--> 275 ancanngtga nagtttncnc gtngangcng aactgtccct gngccnnnac gctccanana 540
W--> 276 cntntccaat ngacaatcga gtttcennnc tccngnaacc tngccgnnnn cnngcccnnc 600
W--> 277 cantntgnta accccgcgcc cggatcgctc tennntcgtt ctncnchnaa ngggntttcn 660
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280 <210> SEQ ID NO: 13
281 <211> LENGTH: 694
282 <212> TYPE: DNA
283 <213> ORGANISM: Homo sapien
285 <220> FEATURE:
286 <221> NAME/KEY: misc_feature
287 <222> LOCATION: (1)...(694)
288 <223> OTHER INFORMATION: n = A,T,C or G
290 <400> SEQUENCE: 13

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FYI

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

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L:23 M:270 C: Current Application Number differs, Wrong Format
L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:45 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:95 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:96 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:119 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:121 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:135 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:142 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:143 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:144 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:145 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:163 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:166 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:169 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:170 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:188 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:189 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
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L:191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
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L:193 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:217 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:269 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:271 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:274 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:275 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:276 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:277 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:299 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:300 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13

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L:301 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:302 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:315 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14